	•
	٠.
	1600/2900
•	CENTER

	CRF Erros Corrected by the STIC Syst ms	rocessing Date:	5 [3
	mb r: 1991925517 Edited	1 by:	(ST
С	Changed a file from non-ASCII to ASCII	d by:	(31
)	Changed the margins in cases where the sequence text was "wrapped" down	n to the next lin	e.
E	Edited a format error in the Current Application Data section, specifically:	ENTE	= K
E a _l	Edited the Current Application Data section with the actual current number.	The number inp	outted by t
A	Added the mandatory heading and subheadings for "Current Application Data	a".	
E	Edited the "Number of Sequences" field. The applicant spelled out a number	instead of usin	g an integ
	Changed the spelling of a mandatory field (the headings or subheadings), spe		
C	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers	that were edite	ed were
In	nserted or corrected a nucleic number at the end of a nucleic line. SEQ ID I	NO's edited:	ā
aj	Corrected subheading placement. All responses must be on the same line as applicant placed a response below the subheading, this was moved to its app	s each subheac propriate place.	ding. If the
lr	Inserted colons after headings/subheadings. Headings edited included:		
	Deleted extra, invalid, headings used by an applicant, specifically:		
- I	Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secreta ☐ page numbers throughout text; ☐ other invalid text, such as	ry initials/filena	me at end
1	Inserted mandatory headings, specifically:		
(Corrected an obvious error in the response, specifically:		
	Edited identifiers where upper case is used but lower case is required, or vice	ce versa.	•
(Corrected an error in the Number of Sequences field, specifically:		
_	A "Hard Page Break" code was inserted by the applicant. All occurrences h	ad to be delete	d.
D dı	Deleted ending stop codon in amino acid sequences and adjusted the *(A)Louise to a Patentin bug). Sequences corrected:	ength:" field ac	cordingly (
	Other:		
,			

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



1642

RAW SEQUENCE LISTING DATE: 05/13/2002 PATENT APPLICATION: US/09/423,351A TIME: 16:42:52

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05132002\I423351A.raw

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SEQUENCE LISTING
      4 (1) GENERAL INFORMATION:
      7
             (i) APPLICANT: The Institute of Cancer Research: Royal
      8
                             Cancer Hospital
     33
            (ii) TITLE OF INVENTION: Binding Complexes
     35
           (iii) NUMBER OF SEQUENCES: 122
     0
            (iv) CORRESPONDENCE ADDRESS:
C-->
     27
                   (B) STREET: 172 Locksley Avenue, Apt #3
     28
                   (C) CITY: San Francisco
                   (D) STATE: California
     29
     30
                   (E) COUNTRY: USA
C--> 31
                   (F) ZIP: CA 94122
C--> 37
             (v) COMPUTER READABLE FORM:
     38
                   (A) MEDIUM TYPE: Floppy disk
     39
                   (B) COMPUTER: IBM PC compatible
     40
                   (C) OPERATING SYSTEM: PC-DOS/MS-DOS
                   (D) SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
     41
C-->43
            (vi) CURRENT APPLICATION DATA:
C--> 44
                   (A) APPLICATION NUMBER: US/09/423,351A
C--> 0
                   (B) FILING DATE: 10-May-2000
C--> 46
           (vii) PRIOR APPLICATION DATA:
                   (A) APPLICATION NUMBER: GB 9710762.7
     47
                   (B) FILING DATE: 23-MAY-1997
     48
C-->
     0
          (viii) ATTORNEY/AGENT INFORMATION:
                   (A) NAME: Liou, Anthony Kian-Fong
     26
     51 (2) INFORMATION FOR SEQ ID NO: 1:
             (i) SEQUENCE CHARACTERISTICS:
     53
     54
                   (A) LENGTH: 15 amino acids
     55
                   (B) TYPE: amino acid
     56
                   (D) TOPOLOGY: linear
     60
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
             Ala Pro Arq Ala Val Phe Pro Ser Ile Val Gly Arg Pro Arg His
     62
                                                                        15
     63
                                                   10
       (2) INFORMATION FOR SEQ ID NO: 2:
     65
             (i) SEQUENCE CHARACTERISTICS:
     67
     68
                   (A) LENGTH: 15 amino acids
     69
                   (B) TYPE: amino acid
     70
                   (D) TOPOLOGY: linear
     74
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
             Phe Pro Ser Ile Val Gly Arg Pro Arg His Gln Gly Val Met Val
     76
     77
             1
                                                   10
                                                                        15
     79 (2) INFORMATION FOR SEQ ID NO: 3:
             (i) SEQUENCE CHARACTERISTICS:
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RAW SEQUENCE LISTING
                                                          DATE: 05/13/2002
                                                           TIME: 16:42:52
                PATENT APPLICATION: US/09/423,351A
                Input Set : A:\PTO.DC.txt
                Output Set: N:\CRF3\05132002\I423351A.raw
82
             (A) LENGTH: 15 amino acids
83
             (B) TYPE: amino acid
84
             (D) TOPOLOGY: linear
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
87
        Gly Arg Pro Arg His Gln Gly Val Met Val Gly Met Gly Gln Lys
89
90
92 (2) INFORMATION FOR SEQ ID NO: 4:
        (i) SEQUENCE CHARACTERISTICS:
94
             (A) LENGTH: 15 amino acids
95
96
             (B) TYPE: amino acid
97
             (D) TOPOLOGY: linear
99
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
         Gly Gly Thr Thr Met Tyr Pro Gly Ile Ala Asp Arg Met Gln Lys
101
                                              10
102
104 (2) INFORMATION FOR SEQ ID NO: 5:
         (i) SEQUENCE CHARACTERISTICS:
106
107
              (A) LENGTH: 15 amino acids
108
              (B) TYPE: amino acid
109
              (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
112
114
         Pro Arg His Gln Gly Val Met Val Gly Met Gly Gln Lys Asp Ser
                          5
                                                                    15
115
         1
                                              10
117 (2) INFORMATION FOR SEQ ID NO: 6:
119
         (i) SEQUENCE CHARACTERISTICS:
120
              (A) LENGTH: 15 amino acids
121
              (B) TYPE: amino acid
122
              (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
125
         Thr Phe Asn Thr Pro Ala Met Tyr Val Ala Ile Gln Ala Val Leu
127
128
                                              10
130 (2) INFORMATION FOR SEQ ID NO: 7:
132
         (i) SEQUENCE CHARACTERISTICS:
133
              (A) LENGTH: 15 amino acids
134
              (B) TYPE: amino acid
135
              (D) TOPOLOGY: linear
138
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
140
         Leu Pro His Ala Ile Leu Arg Leu Asp Leu Ala Gly Arg Asp Leu
141
143 (2) INFORMATION FOR SEQ ID NO: 8:
145
         (i) SEQUENCE CHARACTERISTICS:
146
              (A) LENGTH: 15 amino acids
147
              (B) TYPE: amino acid
              (D) TOPOLOGY: linear
148
152
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
154
        Leu Ala Ser Leu Ser Thr Phe Gln Gln Met Trp Ile Ser Lys Gln
155
                         5
                                              10
         1
157 (2) INFORMATION FOR SEQ ID NO: 9:
159
         (i) SEQUENCE CHARACTERISTICS:
160
              (A) LENGTH: 15 amino acids
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RAW SEQUENCE LISTING DATE: 05/13/2002 PATENT APPLICATION: US/09/423,351A TIME: 16:42:52 Input Set : A:\PTO.DC.txt Output Set: N:\CRF3\05132002\I423351A.raw (B) TYPE: amino acid 161 162 (D) TOPOLOGY: linear 166 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9: 168 Asp Glu Ala Gln Ser Lys Arg Gly Ile Leu Thr Leu Lys Tyr Pro 169 171 (2) INFORMATION FOR SEQ ID NO: 10: (i) SEQUENCE CHARACTERISTICS: 174 (A) LENGTH: 15 amino acids 175 (B) TYPE: amino acid 176 (D) TOPOLOGY: linear 179 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10: Ile Gln Ala Val Leu Ser Leu Tyr Ala Ser Gly Arg Thr Thr Gly 181 182 1 5 10 184 (2) INFORMATION FOR SEQ ID NO: 11: (i) SEQUENCE CHARACTERISTICS: 186 (A) LENGTH: 15 amino acids 187 188 (B) TYPE: amino acid 189 (D) TOPOLOGY: linear 192 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11: 194 Lys Ile Leu Thr Glu Arg Gly Tyr Ser Phe Thr Thr Ala Glu 197 (2) INFORMATION FOR SEQ ID NO: 12: (i) SEQUENCE CHARACTERISTICS: 199 200 -(A) LENGTH: 15 amino acids 201 (B) TYPE: amino acid 202 (D) TOPOLOGY: linear 205 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12: Arg Gly Tyr Ser Phe Thr Thr Ala Glu Arg Glu Ile Val Arg 207 208 1.0 210 (2) INFORMATION FOR SEQ ID NO: 13: 212 (i) SEQUENCE. CHARACTERISTICS: 213 (A) LENGTH: 15 amino acids 214 (B) TYPE: amino acid 215 (D) TOPOLOGY: linear 218 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13: 220 Ala Ser Ser Ser Leu Glu Lys Ser Tyr Glu Leu Pro Asp Gly 5 221 10 15 1 223 (2) INFORMATION FOR SEQ ID NO: 14: (i) SEQUENCE CHARACTERISTICS: 226 (A) LENGTH: 15 amino acids 227 (B) TYPE: amino acid 228 (D) TOPOLOGY: linear 230 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14: 232 Ala Pro Ser Thr Met Lys Ile Lys Ile Ile Ala Pro Pro Glu Arg 233 10 235 (2) INFORMATION FOR SEQ ID NO: 15: 237 (i) SEQUENCE CHARACTERISTICS: 238 (A) LENGTH: 15 amino acids 239 (B) TYPE: amino acid

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DATE: 05/13/2002
                RAW SEQUENCE LISTING
                PATENT APPLICATION: US/09/423,351A
                                                           TIME: 16:42:52
                Input Set : A:\PTO.DC.txt
                Output Set: N:\CRF3\05132002\I423351A.raw
240
              (D) TOPOLOGY: linear
244
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
246
         Ala Pro Pro Glu Arg Lys Tyr Ser Val Trp Ile Gly Gly Ser Ile
247
249 (2) INFORMATION FOR SEQ ID NO: 16:
         (i) SEQUENCE CHARACTERISTICS:
252
              (A) LENGTH: 9 amino acids
253
              (B) TYPE: amino acid
254
              (D) TOPOLOGY: linear
258
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
260
         Ala Val His Ser Gly Ala Leu Asp Asp
                         5
261
         1
263 (2) INFORMATION FOR SEQ ID NO: 17:
         (i) SEQUENCE CHARACTERISTICS:
265
              (A) LENGTH: 9 amino acids
266
267
              (B) TYPE: amino acid
268
              (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
271
273
         Ala Val His Ser Gly Ala Leu Asn Asp
                         5
276 (2) INFORMATION FOR SEQ ID NO: 18:
278
         (i) SEQUENCE CHARACTERISTICS:
279
              (A) LENGTH: 19 amino acids
280
              (B) TYPE: amino acid
281
              (D) TOPOLOGY: linear
283
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
         Ser Gly Ser Gly Gly Arg Pro Arg His Gln Gly Val Met Val Gly Met
285
286
                                              10
288
         Gly Gln Lys
291 (2) INFORMATION FOR SEQ ID NO: 19:
         (i) SEQUENCE CHARACTERISTICS:
293
294
              (A) LENGTH: 19 amino acids
295
              (B) TYPE: amino acid
296
              (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
299
         Ser Gly Ser Gly Ala Arg Pro Arg His Gln Gly Val Met Val Gly Met
301
                         5
                                                                   15
302
         1
                                              10
304
         Gly Gln Lys
307 (2) INFORMATION FOR SEQ ID NO: 20:
309
         (i) SEQUENCE CHARACTERISTICS:
310
              (A) LENGTH: 19 amino acids
311
              (B) TYPE: amino acid
312
              (D) TOPOLOGY: linear
315
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
317
         Ser Gly Ser Gly Gly Ala Pro Arg His Gln Gly Val Met Val Gly Met
318
                                                                   15
320
         Gly Gln Lys
321 (2) INFORMATION FOR SEQ ID NO: 21:
         (i) SEQUENCE CHARACTERISTICS:
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RAW SEQUENCE LISTING
                                                          DATE: 05/13/2002
                PATENT APPLICATION: US/09/423,351A
                                                           TIME: 16:42:52
                Input Set : A:\PTO.DC.txt
                Output Set: N:\CRF3\05132002\I423351A.raw
              (A) LENGTH: 19 amino acids
324
325
              (B) TYPE: amino acid
326
              (D) TOPOLOGY: linear
328
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
         Ser Gly Ser Gly Gly Arg Ala Arg His Gln Gly Val Met Val Gly Met
330
331
                                              10
333
         Gly Gln Lys
336 (2) INFORMATION FOR SEQ ID NO: 22:
         (i) SEQUENCE CHARACTERISTICS:
338
339
              (A) LENGTH: 19 amino acids
340
              (B) TYPE: amino acid
341
              (D) TOPOLOGY: linear
343
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:
345
         Ser Gly Ser Gly Gly Arg Pro Ala His Gln Gly Val Met Val Gly Met
346
                                              10
348
         Gly Gln Lys
351 (2) INFORMATION FOR SEQ ID NO: 23:
         (i) SEQUENCE CHARACTERISTICS:
353
354
              (A) LENGTH: 19 amino acids
355
              (B) TYPE: amino acid
356
              (D) TOPOLOGY: linear
360
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:
         Ser Gly Ser Gly Gly Arg Pro Arg Ala Gln Gly Val Met Val Gly Met
362
363
         1
365
         Gly Gln Lys
367 (2) INFORMATION FOR SEQ ID NO: 24:
369
         (i) SEQUENCE CHARACTERISTICS:
370
              (A) LENGTH: 19 amino acids
371
              (B) TYPE: amino acid
372
              (D) TOPOLOGY: linear
374
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:
376
         Ser Gly Ser Gly Ala Ala Ala Ala Gln Gly Val Met Val Gly Met
377
                                                                   15
                                              10
379
         Gly Gln Lys
382 (2) INFORMATION FOR SEQ ID NO: 25:
         (i) SEQUENCE CHARACTERISTICS:
384
385
              (A) LENGTH: 15 amino acids
386
              (B) TYPE: amino acid
387
              (D) TOPOLOGY: linear
389
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25:
         Met Asp Asp Asp Ile Ala Ala Leu Val Val Asp Asn Gly Ser Gly
391
                          5
                                              10
394 (2) INFORMATION FOR SEQ ID NO: 26:
396
         (i) SEQUENCE CHARACTERISTICS:
397
              (A) LENGTH: 15 amino acids
398
              (B) TYPE: amino acid
399
              (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 26:
401
403
         Ala Ala Leu Val Val Asp Asn Gly Ser Gly Met Cys Lys Ala Gly
```

VERIFICATION SUMMARY

DATE: 05/13/2002

PATENT APPLICATION: US/09/423,351A

TIME: 16:42:53

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05132002\I423351A.raw

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L:7 M:220 C: Keyword misspelled or invalid format, [(i) APPLICANT:] of (1) line corrected
L:12 M:220 C: Keyword misspelled or invalid format, [(F) ZIP:]
L:18 M:220 C: Keyword misspelled or invalid format, [(F) ZIP:]
L:24 M:220 C: Keyword misspelled or invalid format, [(F) ZIP:]
L:31 M:220 C: Keyword misspelled or invalid format, [(F) ZIP:]
L:37 M:220 C: Keyword misspelled or invalid format, [(v) COMPUTER READABLE FORM:]
L:43 M:220 C: Keyword misspelled or invalid format, [(vi) CURRENT APPLICATION DATA:]
L:43 M:220 C: Keyword misspelled or invalid format, [(vi) CURRENT APPLICATION DATA:]
L:44 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:44 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:46 M:220 C: Keyword misspelled or invalid format, [(vii) PRIOR APPLICATION DATA:]
L:0 M:247 C: Inserted Optional Header Field, [(iv) CORRESPONDENCE ADDRESS:]
L:0 M:247 C: Inserted Optional Header Field, [(viii) ATTORNEY/AGENT INFORMATION:]
L:0 M:249 C: Inserted Mandatory Field, [(B) FILING DATE:]
```